WHAT IS CLAIMED IS:

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1	1. A method of sialylating a saccharide group on a recombinant		
2	glycoprotein, the method comprising contacting a saccharide group which comprises	s a	
3	galactose or N-acetylgalactosamine acceptor moiety on a recombinant glycoprotein	with a	
4	sialic acid donor moiety and a recombinant sialyltransferase in a reaction mixture when	nich	
5	provides reactants required for sialyltransferase activity for a sufficient time and unc	er	
6	appropriate conditions to transfer sialic acid from said sialic acid donor moiety to sa	id	
7	saccharide group.		
1	2. The method of claim 1, wherein the sialic acid donor moiety is	CMP-	
2	sialic acid.		
1	 The method of claim 2, wherein the CMP-sialic acid is enzyma 	tically	
2	generated in situ.		
1	4. The method of claim 1, wherein the sialyltransferase is a recom	binant	
2	eukaryotic sialyltransferase which substantially lacks a membrane-spanning domain		
	, , ,		
1	5. The method of claim 1, wherein the sialyltransferase includes a	sialyl	
2	motif which has an amino acid sequence that is at least about 40% identical to a sialyl motif		
3	from a sialyltransferase selected from the group consisting of ST3Gal I, ST6Gal I, and		
4	ST3Gal III.		
I	6. The method of claim 1, wherein the sialyltransferase is a recom	binant	
2	ST3Gal III.		
1	7. The method of claim 6, wherein the sialyltransferase is a recom	hinant	
2	rat ST3Gal III.	Oman	
-	AUT 0 A D CHA AAA.		
1	8. The method of claim 1, wherein the sialyltransferase is a recom	binant	
2	ST3Gal IV.		

Haemophilus 2,3-sialyltransferase.

1		9.	The method of claim 1, wherein the sialyltransferase is a recombinant
2	ST6Gal I.		
1		10.	The method of claim 1, wherein the sialyltransferase is a recombinant
2	ST3Gal I.		
1		11.	The method of claim 10, wherein the reaction mixture comprises a
2	second recom	binan	t sialyltransferase, which second recombinant sialyltransferase is an
3	ST3Gal III.		
1		12.	The method of claim 1, wherein the sialyltransferase is a recombinant
2	bacterial sial	yltrans	ferase.
1		13.	The method of claim 12, wherein the bacterial sialyltransferase has an
2	amino acid sequence which is at least 50% identical to an amino acid sequence of a Neisseria		
3	meningitidis 2,3-sialyltransferase.		
1		14.	The method of claim 13, wherein the bacterial sialyltransferase is a
2	Neisseria meningitidis 2,3-sialyltransferase.		
1		15.	The method of claim 12, wherein the bacterial sialyltransferase has an
2	amino acid s	equen	ce which is at least 50% identical to an amino acid sequence of a
3	Photobacter	ium da	amsela 2,6-sialyltransferase.
1		16.	The method of claim 15, wherein the bacterial sialyltransferase is a
2	Photobacter	ium de	amsela 2,6-sialyltransferase.
		1.7	The south of a Calaine 10 and a sain the heatenial signiferance foreign has an
1		17.	The method of claim 12, wherein the bacterial sialyltransferase has an
2	amino acid sequence which is at least 50% identical to an amino acid sequence of a		

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1	18. The method of claim 17, wherein the sialyltransferase is a Haemophilus		
2	2,3-sialyltransferase.		
1	19. The method of claim 12, wherein the bacterial sialyltransferase has an		
2	amino acid sequence which is at least 50% identical to an amino acid sequence of a		
3	Campylobacter jejuni 2,3-sialyltransferase.		
1	20. The method of claim 19, wherein the sialyltransferase is a		
2	Campylobacter jejuni 2,3-sialyltransferase.		
1	21. The method of claim 1, wherein the sialyltransferase is produced by		
2	recombinant expression of a sialyltransferase in a host cell selected from the group		
3	consisting of an insect cell, a mammalian cell, and a fungal cell.		
	,		
1	22. The method of claim 21, wherein the host cell is an Aspergillus niger		
2	cell.		
1	23. A method of sialylating a saccharide group on a recombinant		
2	glycoprotein, the method comprising contacting a saccharide group which comprises a		
3	galactose or an N-acetylgalactosamine acceptor moiety on a recombinant glycoprotein with a		
4	sialic acid donor moiety and a bacterial sialyltransferase in a reaction mixture which		
5	provides reactants required for sialyltransferase activity for a sufficient time and under		
6	appropriate conditions to transfer sialic acid from said sialic acid donor moiety to said		
7	saccharide group.		
1	 The method of claim 23, wherein the bacterial sialyltransferase has an 		

25. The method of claim 24, wherein the bacterial sialyltransferase is a
 Photobacterium damsela 2,6-sialyltransferase.

amino acid sequence which is at least 50% identical to an amino acid sequence of a

Photobacterium damsela 2,6-sialyltransferase.

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- 1 26. The method of claim 23, wherein the bacterial sialyltransferase has an
 2 amino acid sequence which is at least 50% identical to an amino acid sequence of a *Neisseria*3 *meningitidis* 2,3-sialyltransferase.
- 27. The method of claim 26, wherein the sialyltransferase is a Neisseria
 meningitidis 2,3-sialyltransferase.
- 1 28. The method of claim 23, wherein the bacterial sialyltransferase has an
 2 amino acid sequence which is at least 50% identical to an amino acid sequence of a
 3 Campylobacter jejuni 2,3-sialyltransferase.
 - The method of claim 28, wherein the sialyltransferase is a Campylobacter jejuni 2,3-sialyltransferase.
 - 30. The method of claim 23, wherein the bacterial sialyltransferase has an amino acid sequence which is at least 50% identical to an amino acid sequence of a Haemophilus 2,3-sialyltransferase.
- 31. The method of claim 30, wherein the sialyltransferase is a *Haemophilus* 2.3-sialyltransferase.
 - 32. A method for in vitro sialylation of saccharide groups present on a glycoprotein, said method comprising contacting said saccharide groups with a sialyltransferase, a sialic acid donor moiety, and other reactants required for sialyltransferase activity for a sufficient time and under appropriate conditions to transfer sialic acid from said sialic acid donor moiety to said saccharide group, wherein said sialyltransferase is present at a concentration about 50 mU per mg of glycoprotein or less.
 - 33. The method of claim 32, wherein the sialyltransferase is present at a concentration of between about 5-25 mU per mg of glycoprotein.

1 34. The method of claim 32, wherein the sialyltransferase is present at a 2 concentration of between about 10-50 mU/ml of reaction mixture and the glycoprotein is present in the reaction mixture at a concentration of at least about 2 mg/ml. 3 The method of claim 32, wherein the method yields a glycoprotein 1 2 having sialylation of at least about 80% of terminal galactose residues present on the 3 saccharide groups. The method of claim 32, wherein the sialyltransferase is a recombinant 1 36. 2 sialyltransferase. The method of claim 36, wherein the sialyltransferase substantially 1 2 lacks a membrane-spanning domain. 38. The method of claim 32, wherein the sialyltransferase includes a sialyl 1 motif which has an amino acid sequence that is at least about 40% identical to a sialyl motif 2 3 from a sialyltransferase selected from the group consisting of ST3Gal I, ST6Gal I, and 4 ST3Gal III. The method of claim 32, wherein the sialyltransferase is an ST3Gal III. 1 1 40. The method of claim 39, wherein the ST3Gal III is a rat ST3Gal III. The method of claim 32, wherein the sialyltransferase is an ST3Gal IV. 1 1 The method of claim 32, wherein the sialyltransferase is an ST3Gal I. 42. 1 The method of claim 42, wherein the reaction mixture comprises a second recombinant sialyltransferase, which second recombinant sialyltransferase is an 2 3 ST3Gal III.

- 44. The method of claim 32, wherein the sialyltransferase is a bacterial 1 2 sialvltransferase. 1 The method of claim 44, wherein the bacterial sialyltransferase is a 2 recombinant sialyltransferase. 1 46. The method of claim 44, wherein the bacterial sialyltransferase has an amino acid sequence which is at least 50% identical to an amino acid sequence of a Neisseria 2 3 meningitidis 2,3-sialyltransferase. 1 47. The method of claim 46, wherein the bacterial sialyltransferase is a 2 Neisseria meningitidis 2,3-sialyltransferase. 1 48. The method of claim 44, wherein the bacterial sialyltransferase has an amino acid sequence which is at least 50% identical to an amino acid sequence of a 3 Photobacterium damsela 2,6-sialyltransferase. 1 49. The method of claim 48, wherein the bacterial sialyltransferase is a Photobacterium damsela 2.6-sialvltransferase. 1 50. The method of claim 44, wherein the bacterial sialyltransferase has an 2 amino acid sequence which is at least 50% identical to an amino acid sequence of a 3 Campylobacter jejuni 2,3-sialyltransferase. 1 51. The method of claim 50, wherein the sialyltransferase is a Campylobacter jejuni 2,3-sialyltransferase.
- 1 52. The method of claim 44, wherein the bacterial sialyltransferase has an
 2 amino acid sequence which is at least 50% identical to an amino acid sequence of a
 3 *Haemophilus* 2,3-sialyltransferase.

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- 1 53. The method of claim 52, wherein the sialyltransferase is a *Haemophilus*2 2,3-sialyltransferase.

 1 54. The method of claim 32, wherein the sialic acid donor moiety is CMP2 sialic acid.

 1 55. The method of claim 54, wherein the CMP-sialic acid is enzymatically
 2 generated *in situ*.
 - 56. The method of claim 32, wherein the sialic acid is selected from the group consisting of NeuAc and NeuGc.
 - 57. A method for in vitro sialylation of saccharide groups present on a glycoprotein, the method comprising contacting the saccharide groups with an ST3Gal III sialyltransferase, a sialic acid donor moiety, and other reactants required for sialyltransferase activity for a sufficient time and under conditions to transfer sialic acid from said sialic acid donor moiety to said saccharide group, wherein said ST3Gal III sialyltransferase is present at a concentration of less than about 50 mU per mg of glycoprotein.
- 1 58. The method of claim 57, wherein the method further comprises 2 contacting the saccharide groups with an ST6GalI sialyltransferase.